Figure 1

	10	24	20	40	£ A
	-	20	30	40	50
CATGAC	CTCC	actguaagag	GGGGCTAGCG	TGAGCGCTGA	TTCTCAACCT
	60	70	80	90	100
ACCATA	ACTC	TTTCCTGCCT	CAGGAACTCC	AATAAAACAT	TTTCCATCCA
102					
AC	3'				

Figure 3

ATCTACCAGCTCATGATGCAGTGCTGGCAGCAGGAGCGTGCCCACCGCCC
CAAGTTCGCTGACATCGTCAGCATCCTGGACAAGCTCATTCGTGCCCCTG
ACTCCCTCAAGACCCTGGCTGACTTTGACCCCCGCGTGTCTATCCGGCTC
CCCAGCACGAGCGGCTCCGGAGGGGGGTGCCCTTCCGCACGGTGTCCGAGT
GGCTGGAGTCCATCAAGATGCAGCAGTATACGGAGCACTTC

Figure 4

CATGCATCACGGATCAATAGACTGTACTTATTTTCCAATAAAATTTTCAA ACTTTGTACTGTT

Figure 6

AACTTGCCCTGTGCCTGTGTCCCCCATGCTAGGGGCGGAGGGGTCTTTTC
CTTCTTCTTTCCTACCTACCCCTTTTCTCTTTGGCCAGGGGCCTCGTATCCT
ACCTTTCCTTGTCCCCTGGGCTGCACAGAGGATTGCCCCTTCTCTTT
TCAGAGCTGGCCCTCGATGCCAAATTAGCATTTAGTATTTTGCTCAAAGTC
TAAGGGACC

Figure 7

Figure 9

Figure 10

Figure 12

Figure 13

CATGGAGCAGCCCCTGTTCCGGGGGCAGCCAGTGACCCAGCCCACC
AATGGGCCTCCAGGACCCCAGGAACAATAAAATGTCTTCTCCCACC

Figure 15

Figure 16

Figure 18

Figure 19

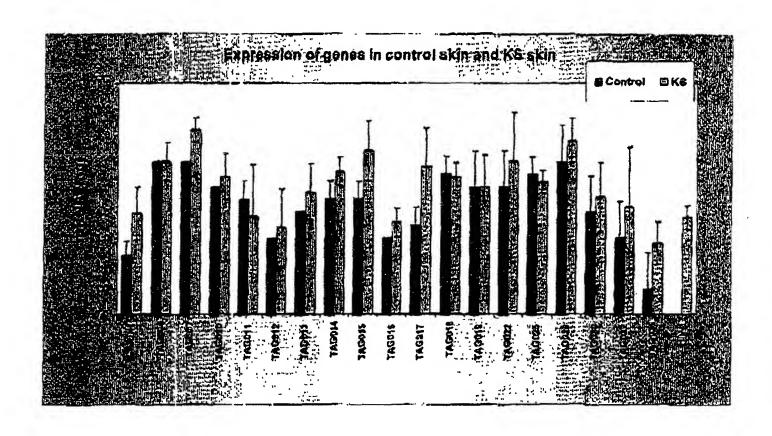


Figure 20

